

Table S1: List of primers.

Cloning of <i>znf385c</i> full-length and splice variants	ct39a_GSP_R	TTCCTTCGATGGCGGTGTCGTCCAG
	FT39_fl_5'_F	GGAAACAAGGATAAGCCTACC
	FT39a_fl_F	ATaAGCCTACCTTTTCACAA
	FT39a_v2_R	TTGCCTGCAGTATCTTGTCTTT
	FT39a_v1_R	GGGTAGATCAATAGGGGGAGA
In situ hybridization	<i>v1_ish_F</i>	TAATGATGGATGGACAGAGTGTG,
	<i>v1_ish_R</i>	GGGGGTAGATCAATAGGGGGAGA;
	<i>v2_ish_F</i>	TGTAGCCTTTTTCTAGACCTGTTT,
	<i>v2_ish_R</i>	GCCGTGCGGTATCAAACATT
Gene expression profile by quantitative PCR	p21_QPCR_F	AAACATCCCGAAAACACCAG
	p21_QCPR_R	ACGCTTCTTGGCTTGGTAGA;
	mdm2_QPCR_F	CAGGAGGAGGAGAAGCAGTG
	mdm2_QPCR2_R	GCTGTCCGACTTATGCCTCT
	bclxL_QPCR_F	GATCCAAAGCCAAGGAGGAT
	bclxL_QPCR_R	TCTTGCGATTTCTTGCTTTC;
	p53_QPCR_F	ttccggagacaagcgactat
	p53_QPCR_R	caggtccggtgaataagtgc;
	puma_QPCR_F	ATGATGCCTTCAGCTTGGAC
	puma_QPCR_R	CCTCTGAGGAGCTCTGGTTG;
	noxal_QPCR_F	ATGGCGAAGAAAGAGCAAAC
	noxal_QPCR_R	CGCTTCCCCTCCATTGTGA.
	beta actin_F:	AATCCCAAAGCCAACAGAGA,
	beta actin_R	ACATACATGGCAGGGGTGTT.
Chromatin immunoprecipitation (ChIP)	<i>p21^{cip1/waf1}</i> +1kb	F: tcagggtgtggtgtggttaa R: aatgggcaacatgttcttag
	<i>p21^{cip1/waf1}</i> +1.56kb	F: gtcattggactacagtggttga R: agggagcagcgctagacat
	<i>p21^{cip1/waf1}</i> +1.7kb	F: tetgtgactcactgcccatt, gcgcctctgagctgtcat
	<i>p21^{cip1/waf1}</i> +2.2kb	F: gcaaaaccctatttttagcatgg R: gcaatgtgagctaacaacaagag).
	<i>mdm2</i> -2kb	F: actcgcattccaagtcttg R: aagatccacgcaaaaggatg),
	<i>mdm2</i> -1.5kb	F: aacgcattcatgcctgttta, R: aaatcagcatgactcagaaggat
	<i>mdm2</i> -1.1kb	F: cccaaacatgttgagcgtag, R: aggtcttttagggccagggtg